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Our Case No. 5404/18

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Application of:)
Atsuhiko Shinmyo et al.)
International Application No. PCT/JP01/05096)
International Filing Date: June 15, 2001)
Title of Invention: METHOD OF INDUCING GENE EXPRESSION IN PLANT AND PLANT TREATED)
THEREBY	"

PRELIMINARY AMENDMENT

Attn: Box DO/EO/US Commissioner for Patents Washington, D.C. 20231

Dear Sir:

Preliminary to examination of the subject application, please amend the aboveidentified application as follows:

IN THE SPECIFICATION

Page 1, after the title, insert the following paragraph:

"RELATED APPLICATIONS

This application is a nationalization of PCT application PCT/JP01/05096 filed June 15, 2001. This application claims priority from the PCT application and Japan Application Serial No. 2000-180466 filed June 15, 2000.

IN THE CLAIMS

Please amend the claims as follows:

- 4. (Amended) The method according to Claim 1, wherein said autogenous regulatory factor is a butyrolactone autogenous regulatory factor.
- 5. (Amended) The method according to Claim 1, wherein said autogenous regulatory factor is virginiae butanolide.
- 6. (Amended) The method according to Claim 1, wherein said gene expression inducing system is involved in a production of an antibiotic.
- 7. (Amended) The method according to Claim 1, wherein said gene expression inducing system is involved in a production of virginiamycin.
- 8. (Amended) The method according to Claim 1, wherein said repressor gene is a barA gene.
- 9. (Amended) The method according to Claim 1, wherein said repressor gene contains a region comprising a nucleotide sequence shown under SEQ ID NO:1.
- 10. (Amended) The method according to Claim 1, wherein said repressor gene contains a region coding for an amino acid sequence shown under SEQ ID NO:2.
- 11. (Amended) The method according to Claim 1, wherein a promoter for said repressor gene is a plant promotor.

- 13. (Amended) The method according to Claim 1, wherein a nucleotide sequence of said operator is derived from a barA, barB or barX gene.
- 14. (Amended) The method according to Claim 1, wherein a nucleotide sequence of said operator is BARE-1, BARE-2 or BARE-3.
- 15. (Amended) The method according to Claim 1, wherein a nucleotide sequence of said operator is BARE-3.
- 16. (Amended) The method according to Claim 1, wherein the nucleotide sequence of said operator contains a region comprising a nucleotide sequence shown under SEQ ID NO:3.
- 17. (Amended) The method according to Claim 1, wherein a promoter for said gene placed under the control of the operator is a plant promoter.
- 19. (Amended) The method according to Claim 17, wherein said operator is disposed in at least one place in said plant promoter.
- 20. (Amended) The method according to Claim 17, wherein said operator is disposed in at least one place in the vicinity of a site 3' downstream or in the vicinity of a site 5' upstream of a TATA box of said plant promoter.

- 21. (Amended) The method according to Claim 17, wherein said operator is disposed, together with the TATA box of said plant promoter, in a manner shown under any of SEO ID NO:4 through SEQ ID NO:7.
- 22. (Amended) The method according to Claim 1, wherein said gene placed under the control of the operator is a gene capable of providing the plant with fertility.
 - 23. (Amended) A plant transformed by the method according to Claim 1.
- 24. (Amended) Tobacco (<u>Nicotiana tabacum</u> L.) transformed by the method according to Claim 1.
- 25. (Amended) A cultured plant cell transformed by the method according to Claim 1.
- 26. (Amended) A cultured tobacco cell transformed by the method according to Claim 1.
- 27. (Amended) A cultured tobacco BY2 cell transformed by the method according to Claim 1.
- 30. (Amended) The repressor gene according to Claim 28 wherein said repressor gene is a barA gene.

- 31. (Amended) The repressor gene according to Claim 28 wherein said repressor gene contains a region comprising a nucleotide sequence shown under SEQ ID NO:1.
- 32. (Amended) The repressor gene according to Claim 28 wherein said repressor gene contains a region coding for an amino acid sequence shown under SEQ ID NO:2.
- 35. (Amended) The modified promoter according to Claim 33, wherein a nucleotide sequence of said operator is BARE-1, BARE-2 or BARE-3.
- 36. (Amended) The modified promoter according to Claim 33, wherein the nucleotide sequence of said operator contains a region comprising a nucleotide sequence shown under SEQ ID NO:3.
- 37. (Amended) The modified promoter according to Claim 33, wherein said operator is disposed, together with the TATA box of said plant promoter, in a manner shown under any of SEQ ID NO:4 through SEQ ID NO:7.

REMARKS

The specification is amended to make record therein of the priorities claimed. The Claims are amended to reduce their number.

Respectfully submitted,

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Reg. No. 19,795

Attorney for Applicants

Date:

Feb. 15,2002

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APPENDIX A

Version With Markings to Show Changes Made

CLAIMS

- 4. (Amended) The method according to [any of] Claim[s] 1 [to 3], wherein said autogenous regulatory factor is a butyrolactone autogenous regulatory factor.
- 5. (Amended) The method according to [any of] Claim[s] 1 [to 3], wherein said autogenous regulatory factor is virginiae butanolide.
- 6. (Amended) The method according to [any of] Claim[s] 1 [to 5], wherein said gene expression inducing system is involved in a production of an antibiotic.
- 7. (Amended) The method according to [any of] Claim[s] 1 [to 5], wherein said gene expression inducing system is involved in a production of virginiamycin.
- 8. (Amended) The method according to [any of] Claim[s] 1 [to 7], wherein said repressor gene is a barA gene.
- 9. (Amended) The method according to [any of] Claim[s] 1 [to 8], wherein said repressor gene contains a region comprising a nucleotide sequence shown under SEQ ID NO:1.

- 10. (Amended) The method according to [any of] Claim[s] 1 [to 9], wherein said repressor gene contains a region coding for an amino acid sequence shown under SEQ ID NO:2.
- 11. (Amended) The method according to [any of] Claim[s] 1 [to 10], wherein a promoter for said repressor gene is a plant promotor.
- 13. (Amended) The method according to [any of] Claim[s] 1 [to 12], wherein a nucleotide sequence of said operator is derived from a barA, barB or barX gene.
- 14. (Amended) The method according to [any of] Claim[s] 1 [to 12], wherein a nucleotide sequence of said operator is BARE-1, BARE-2 or BARE-3.
- 15. (Amended) The method according to [any of] Claim[s] 1 [to 12], wherein a nucleotide sequence of said operator is BARE-3.
- 16. (Amended) The method according to [any of] Claim[s] 1 [to 15], wherein the nucleotide sequence of said operator contains a region comprising a nucleotide sequence shown under SEQ ID NO:3.
- 17. (Amended) The method according to [any of] Claim[s] 1 [to 16], wherein a promoter for said gene placed under the control of the operator is a plant promoter.

- 19. (Amended) The method according to Claim 17 [or 18], wherein said operator is disposed in at least one place in said plant promoter.
- 20. (Amended) The method according to Claim 17 [or 18], wherein said operator is disposed in at least one place in the vicinity of a site 3' downstream or in the vicinity of a site 5' upstream of a TATA box of said plant promoter.
- 21. (Amended) The method according to [any of] Claim[s] 17 [to 20], wherein said operator is disposed, together with the TATA box of said plant promoter, in a manner shown under any of SEQ ID NO:4 through SEQ ID NO:7.
- 22. (Amended) The method according to [any of] Claim[s] 1 [to 21], wherein said gene placed under the control of the operator is a gene capable of providing the plant with fertility.
- 23. (Amended) A plant transformed by the method according to [any of] Claim[s] 1 [to 22].
- 24. (Amended) Tobacco (Nicotiana tabacum L.) transformed by the method according to [any of] Claim[s] 1 [to 22].
- 25. (Amended) A cultured plant cell transformed by the method according to [any of] Claim[s] 1 [to 22].

- 26. (Amended) A cultured tobacco cell transformed by the method according to [any of] Claim[s] 1 [to 22].
- 27. (Amended) A cultured tobacco BY2 cell transformed by the method according to [any of] Claim[s] 1 [to 22].
- 30. (Amended) The repressor gene according to Claim 28 [or 29] wherein said repressor gene is a barA gene.
- 31. (Amended) The repressor gene according to [any of] Claim[s] 28 [to 30] wherein said repressor gene contains a region comprising a nucleotide sequence shown under SEQ ID NO:1.
- 32. (Amended) The repressor gene according to [any of] Claim[s] 28 [to 31] wherein said repressor gene contains a region coding for an amino acid sequence shown under SEQ ID NO:2.
- 35. (Amended) The modified promoter according to Claim 33 [or 34], wherein a nucleotide sequence of said operator is BARE-1, BARE-2 or BARE-3.
- 36. (Amended) The modified promoter according to [any of] Claim[s] 33 [to 35], wherein the nucleotide sequence of said operator contains a region comprising a nucleotide sequence shown under SEQ ID NO:3.

37. (Amended) The modified promoter according to [any of] Claim[s] 33 [to 36], wherein said operator is disposed, together with the TATA box of said plant promoter, in a manner shown under any of SEQ ID NO:4 through SEQ ID NO:7.



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<301> Okamoto, S., Nakamura, K., Nihira, T. and Yamada, Y.

<302> Virginiae butanolide binding protein from Streptomyces virginiae. Evidence that VbrA is not the virginiae butanolide binding protein and reidentification of the true binding protein

<303> The Journal of Biological Chemistry

<304> 270

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 <301> Kinoshita, H., Tsuji, T., Ipposhi, H., Nihira, T. and Yamada, Y.
 <302> Characterization of Binding Sequences for Butyrolactone Autoregulator
 Receptors in Streptomycetes
 <303> Journal of Bacteriology
 <304> 181
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